

20 30 40 50 60  
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 TCTCTTCCTG GATCCTTCAG AGCTCTTGTG AATTCCAC GTTTTTTTTT GTTTTTTCGT  
 130 140 150 160 170 180  
 CGTTTAATTG TGGAAACACA TATCCGTCCCT CTTTGAAACA GCATCAGAAA ACTTTCTGCT  
 190 200 210 220 230 240  
 CTCCGTGTCC TTCTACTTAC TCTGATTGCC TTAGTTAGTC ACATCGCAAG CAACAACCAA  
 250 260 270 280 290 300  
 CTGCCAATGG GAGGAGCCAG TTGGAGCAGG GTGCGTGCTC GGTGCTCTTT TCAGAAGGTT  
 310 320 330 340 350 360  
 TTCTCTTGTG CCAGCATGCT TTTTGAGGC TGTGTCATCA CAATGAACAT GTGTGAGTTC  
 370 380 390 400 410 420  
 ATCCGTCTGG ATTATTCTTT TTCTTACGTC TTCTGAGTAC TTCATACTTT CCAAATTTTT  
 430 440 450 460 470 480  
 CAACTGAACT TTTCTTCTTT TCTCATTGAA GTGGTTGGT TTTGGTCGCG TGATCAACGG  
 490 500 510 520 530 540  
 ATCCTACTTT TTTGAACAA AATGTTTTG AAGTTTCACA GACTGATTTTC GGGGTTTTTT  
 550 560 570 580 590 600  
 CAAAGAATAT ATTCCCTCTC GAGCAAGAGA AAATTCCAGA AAATAGTAGT TTTTTTCAAT  
 610 620 630 640 650 660  
 TAGTCGTTTC ATTTGTACTA GCTAAAAC TTGCAACTTA TGGCTTTAAA ACATGTGTTG  
 670 680 690 700 710 720  
 GCTTCATACA AAAACATTAA ACTAGTGTGTT TTCCAGTTTT GTGTTCGTTT CATTTCCTCA  
 730 740 750 760 770 780  
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 TTTCTCTTCT CACGTCTTGT TATTTCTCG ATTTTATTTT CTGAATCTGT GCGGTTTCA  
 850 860 870 880 890 900  
 ATCAATTGAA TTGCGATAAT TATTCTATCA GAAATATATT TTCAGAAATC CAAATACTCC  
 910 920 930 940 950 960  
 AGGTGCCAAT GCGGTGAAAG AAAATTC ~~TGA~~ AGTTTATTCC TGAAATCACA CTACTCTTGC  
 970 980 990 1000 1010 1020  
 TTTTATTTGT ACACCTACAA CAGGTTAGTT GGTTGATTCT AGATCTCTTG CCTCCTAGCT  
 1030 1040 1050 1060 1070 1080  
 TGCAAGGATA ATATAATTGA ATTGTTTTG AGGAGTGCAA AGATTGAATA GTTTCTATA  
 1090 1100 1110 1120 1130 1140  
 TTTAGGCTAA AGGAAACGA CGGAAATGTC CGGAGGGTGC GTGGTCGGAA GGAAAGATTA

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GTCTGAACTT TTGATTCAA ACTTTCCGAA CTTCAAGCG GTTCCAAATT ACTCACTTCC					
1390	1400	1410	1420	1430	1440
ATTTATCTCT TTGCTACAAT TTCTCCCACA AAGCCTTTT CTTCATTAA CGTTCTTTT					
1450	1460	1470	1480	1490	1500
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1510	1520	1530	1540	1550	1560
TCACATGAGG ATAAATTGAA TGGAATGACA AGTTTGTGC CCAGAAGGCA GTTTGCACT					
1570	1580	1590	1600	1610	1620
GAACCTGTTG AGTTGCAGAC ACATCTCAA ACACAGAAGA TGAGTGGAAA ACTAGTGAGA					
1630	1640	1650	1660	1670	1680
GACTGCCAAA AGTCGAAGGG ATAATGAAAA TTTCTGCAA ATGAATTCTG CGAAGTTATG					
1690	1700	1710	1720	1730	1740
TGAAAAATTAA TTGGATTGGG AGTTGTGGG GTGAAGAGAT GGGTCAAAG CCATCAATCT					
1750	1760	1770	1780	1790	1800
TGAATGCTTC GGTCAAAGAT TTGTTCTCA TATGTTACA ACACTGAAAA CAATCTATCC					
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TAGAAATGTT TGAACCACCC TCTAAAGTCC TTCCGTATAT TTTTCATCT TTATACCGAC					
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CAAGTTGGCG GCACCTTGGA GAGTGAATTG GATGAAAAAG TGTTTGATAA GTTTGTCGGG					
2050	2060	2070	2080	2090	2100
CAAACGTGTC CCCTGGGTGG GGAAATGGTG GCATTTTTGG AAACATTTTC ATAGTCGAAG					
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AAGTGGAAACA AGAAATTGGA AAAATAGAGA TACATATGTA TATGAAAATA GAATTGAAACA					
2170	2180	2190	2200	2210	2220
GGAACCTTATT TTTATTTCA GGATATGGGA AGCTTGAATG AAATATCATE CGACTTTGAA					
2230	2240	2250	2260	2270	2280
ATTGACATT TATTCACTCA ACTGTGGCAT GACTCGGCAC TTTCTTTGC TCATCTTCCG					

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 2350 2360 2370 2380 2390 2400  
 TCAGAAATAT CACAATGGAA ACACGACTTT TACCTAAGAT TTGGTCTCCA AACACGTGTA  
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 TGATTAATTC AAAACGAACA ACCGTCCATG CATCACCACATC GGAAAATGTG ATGGTTATTTC  
 2470 2480 2490 2500 2510 2520  
 TGTACGAGGT ATGATTTTG ATTTTGTGAC GTCACAAACA GAGCATGTCT AAGGGCATGT  
 2530 2540 2550 2560 2570 2580  
 TGTAGCAAGA AAAAACCGGA TTCTTGTCTC TGTCGACGTT TCCTAAGTAT TGTGAATTAT  
 2590 2600 2610 2620 2630 2640  
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 2830 2840 2850 2860 2870 2880  
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 2950 2960 2970 2980 2990 3000  
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 3010 3020 3030 3040 3050 3060  
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 3130 3140 3150 3160 3170 3180  
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 3190 3200 3210 3220 3230 3240  
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 3250 3260 3270 3280 3290 3300  
 AACCAAAAGC TCTGCGGGCA AGAACAACTG TCGGAATCTC ATCTCTTCTA GCTCTTACTT  
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 3370 3380 3390 3400 3410 3420  
 TTTTCTTTTT CAACAAATA AAAAAGA TAAACAAATA TTTGTTTCAG CAATGGATGT

3410	3440	3450	3460	3470	3480
GTGGATGCTT GGATGCATAT CATTGTCCTT CGGAACCAGT GTAGAATTGG CATTGTTTG					
3490	3500	3510	3520	3530	3540
TTACATTTCG CGTTGTCAGA ACAGCGTAAG AAAGTGAGTT GGCATAAGAG TTTTCTCACG					
3550	3560	3570	3580	3590	3600
TGGAGGGAAG TAATTAATT TTGGGTGTCA TATGAAAATA TCAAAACAA TATCAGGAAA					
3610	3620	3630	3640	3650	3660
TTGAATTCA CTATGATTTC CTAGTAAACA AATTACAGCG CGGAACGACG ACGGAAACGA					
3670	3680	3690	3700	3710	3720
ATGAGAAATT CTCAGGTGTG GGCAAACGGA TCGTCTAGAA CTAGAAGCAA CGGGTATGCA					
3730	3740	3750	3760	3770	3780
AACGGGGAT CTGTAATCTC ACATTATCAT CCAACAAGCA ATGGAAATGG GAATAATAAT					
3790	3800	3810	3820	3830	3840
CGACATGATA CACCTCAAGT TACTGGAAGG TTAGCAATCT CTATGATAGC ATTTATCAAT					
3850	3860	3870	3880	3890	3900
TATTAAAGAA CTCTGGAATT AGTTTTAAA GTATAAATAA ATCTCTATT CTTGCGACCT					
3910	3920	3930	3940	3950	3960
ACATTGAACT TAATAGTTAT GTTTTACAGA GGATCACTTC ATCGAAACGG GCCACCAC					
3970	3980	3990	4000	4010	4020
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4030	4040	4050	4060	4070	4080
AGTCTTACAT TGAGTTCAAA CTTTTGAAAT TTAAGCGTTC TATCTGATAA AGTTCTTCGG					
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TGGTTTATA ATTTTGATT CATAAACTTA CCCACTCCTT TCTCACTAAC ATTTTACCT					
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4210	4220	4230	4240	4250	4260
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4270	4280	4290	4300	4310	4320
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4330	4340	4350	4360	4370	4380
GTTATGGAAA TCACATATAC TTTGTTCTGG AATTGTATAT GTATGCTTTG AAAAAGCACA					
4390	4400	4410	4420	4430	4440
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4450	4460	4470	4480	4490	4500
GACACTCTTA AGTTATCATA TTCTAATTTC CAAGAATGTT ATATTTGAA GAAGCCGGTG					
4510	4520	4530	4540	4550	4560
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 4630 4640 4650 4660 4670 4680  
 TCTGTCTGAT CATGGTTTTC GGACTGAAAT TGTGTTAAC TGAAGTTATA TGTGAGCCAC  
 4690 4700 4710 4720 4730 4740  
 ATTGATTAAA CCTGTGAGAG ATGCCCATTT GTACTCATT TACGACTGTC TCATGTCAA  
 4750 4760 4770 4780 4790 4800  
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 CCACCATCTG GATGTCTGGC CAGATTCCAT CCGGAAGCAG TGGACAAATT CTCCATTGTA  
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 4930 4940 4950 4960 4970 4980  
 AATCATAAAAT ATCTCGACTT TTCAGCTTGT CTACTGGTGG CACTATTGTT CTCAAACTTT  
 4990 5000 5010 5020 5030 5040  
 CGATCAAAAC TATCAGTGAT TGAAGTTTAT CCCTTTAAT TCCAATAATT CACAGTTGCC  
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 5170 5180 5190 5200 5210 5220  
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 5350 5360 5370 5380 5390 5400  
 TTATATTGCT CGCACCCCTAA ATGACAGGTA TTAGAAATTA ACCGCTTTTC AGAGTATT  
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 5470 5480 5490 5500 5510 5520  
 CTCAACTTCA GTCGGACAAA TTTTAAATT TTTACTCGAT AAAAAAATT TATAATTGAG  
 5530 5540 5550  
 ACAAAATTATG TCTTCTCATT TTTGATCGCT

Fig. 1  
Page 5

20 30 40 50  
 ATGAAGTTAA TTCCTGAAAT CACACTACTC TTGCTTTAT TGTACACTC  
 60 70 80 90 100  
 TACACAGGCT AAAGGAAAAC GACGGAAATG TCCGGAGGGT GCGTGGTCGG  
 110 120 130 140 150  
 AAGGAAAGAT TATGAAACACG ATCATGAGCA ACTACACGAA AATGTTGCC  
 160 170 180 190 200  
 GACGCGGAGG ACAGCGTACA AGTTAATATT GAGATTCATG TACAGGATAT  
 210 220 230 240 250  
 GGGAGCTTG AATGAAATAT CATCCGACTT TGAAATTGAC ATTTTATTCA  
 260 270 280 290 300  
 CTCAACTGTG GCATGACTCG GCACTTTCTT TTGCTCATCT TCCGGCTTGT  
 310 320 330 340 350  
 AAGCGAAATA TCACAAATGGA AACACGACTT TTACCTAAGA TTTGGTCTCC  
 360 370 380 390 400  
 AACACACGTGT ATGATTAATT CAAAACGAAC AACCGTCCAT GCATCACCAT  
 410 420 430 440 450  
 CGGAAAATGT GATGGTTATT CTGTACGAGA ATGGAACAGT CTGGATTAAC  
 460 470 480 490 500  
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 TTTCGATACT CAAACTTGCA TATTAATCTT TGAATCCTAT AGTCATAACT  
 560 570 580 590 600  
 CAGAAGAAGT TGAACTTCAT TGGATGGAAG AAGCTGTCAC ATTAATGAAG  
 610 620 630 640 650  
 CCAATTCAAC TTCCCTGACTT TGATATGGTT CATTATTCAA CTAAAAGGA  
 660 670 680 690 700  
 AACTTTACTC TATCCAAACG GGTACTGGGA TCAGCTTCAA GTTACTTTCA  
 710 720 730 740 750  
 CTTTCAAACG ACGATATGGA TTCTATATTA TTCAAGCCTA TGTTCCAACA  
 760 770 780 790 800  
 TATCTTACAA TCATTGTATC TTGGGTTTCA TTCTGCATGG AACCAAAGC  
 810 820 830 840 850  
 TCTGCCGGCA AGAACAACTG TCGGAATCTC ATCTCTTCTA GCTCTTACTT  
 860 870 880 890 900  
 TCCAGTTGG AAAATTTTG AAAAATCTTC CAAGGGTTTC ATATGTGAAA  
 910 920 930 940 950  
 GCAATGGATG TGTGGATGCT TGGATGCATA TCATTTGTCT TCGGAACCAT

970 980 990 1000  
GGTAGAA~~T~~ GCATTTGTTT GTTACATTT~~C~~ CCGTTGTCAC~~T~~ CAGCGTAA

1010 1020 1030 1040 1050  
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1060 1070 1080 1090 1100  
AACGGATCGT GTAGA~~T~~ACTAG AAGCAACGGG TATGCAAACG GGGGATCTGT

1110 1120 1130 1140 1150  
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1160 1170 1180 1190 1200  
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1210 1220 1230 1240 1250  
CCATCTCCAT TAAACCTTCA AATGACTACA TTTGATT~~CG~~ AGATCCCTCT

1260 1270 1280 1290 1300  
GACTTTGAT CAGCTGCCAG TTTCCATGGA ATCCGATAGA CCCCTGATTG

1310 1320 1330 1340 1350  
AAGAGATGCG ATCAACATCA CCACCTCCAC CATCTGGATG TCTGGCCAGA

1360 1370 1380 1390 1400  
TTCCATCCGG AAGCAGTGG~~A~~ CAAATTCTCC ATTGTAGCTT TTCCATTGGC

1410 1420 1430 1440 1450  
ATTTACAATG TTTAATCTTG TCTACTGGTG GCACTATTTG TCTCAAAC~~T~~

1460 1470  
TCGATCA~~AAA~~ CTATCAGTGA

Fig. 2  
Page 2

20 30 40 50  
MKF IPEIY LLLFVHSTQA KGKRRKCPEG AWSEGKIMN SNYTKMLP  
60 70 80 90 100  
DAEDSVQVN EIHVQDMGSL NEISSLDFEID ILFTQLWHDS ALSFAHLPAC  
110 120 130 140 150  
KRNITMETRL LPKIWSPNTC MINSKRTTVH ASPSENVMVI LYENGTVWIN  
160 170 180 190 200  
HRLSVKSPCN LDLRQFPFDT QTCILIFESY SHNSEEVELH WMEEAVTLMK  
210 220 230 240 250  
PIQLPDFDMV HYSTKKETLL YPNGYWDQLQ VTFTFKRRYG FYIIQAYVPT  
260 270 280 290 300  
YLTIIVSWVS FCMEPKALPA RTTVGISSLL ALTFQFGNIL KNLPRVSYVK  
310 320 330 340 350  
AMDVWMLGCI SFVFGTMVEL AFVCYISRCQ NSVRNAERRR ERMRNSQVWA  
360 370 380 390 400  
NGSCRTRSNG YANGGSVISH YHPTSNGNGN NNRHDTPQVT GRGSLHRNGP  
410 420 430 440 450  
PSPLNLQM TT FDSEIPLTFD QLPVSMESDR PLIEEMRSTS PPPPSGCLAR  
460 470 480  
FHPEAVDKFS IVAFFLAFTM FNLVYWWHYL SQTFDQNYQ

Fig. 3

● MOD-1 is similar  
to ligand-gated ion channels

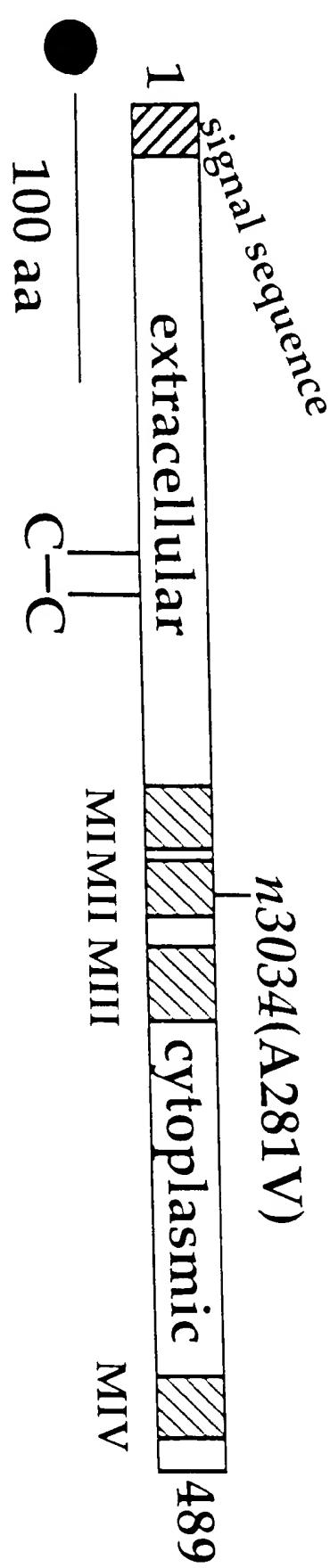


Fig. 4

ok103 is a 4135 bp deletion  
allele of *mod-1*

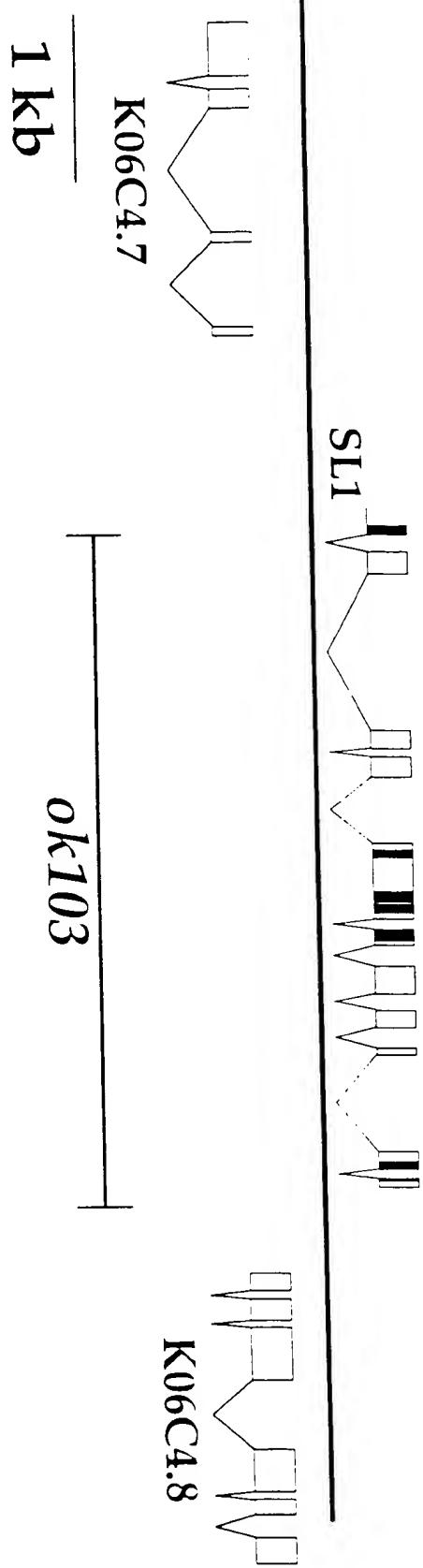


Fig. 5

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 130 140 150 160 170 180  
 CGTTTAATTG TGGAAACACA TATCCGTCCT CTTTGAAACA GCATCAGAAA ACTTTCTGCT  
 190 200 210 220 230 240  
 CTCCGTGTCC TTCTACTTAC TCTGATTGCC TTAGTTAGTC ACATCGCAAG CAACAACATAA  
 250 260 270 280 290 300  
 CTGCCAATGG GAGGAGCCAG TTGGAGCAGG GTGCGTGCTC GGTGCTCTTT TCAGAAGGTT  
 310 320 330 340 350 360  
 TTCTCTTGTG CCAGCATGCT TTTTGAGGC TGTGTCATCA CAATGAACAT GTGTGAGTTC  
 370 380 390 400 410 420  
 ATCCGTCTGG ATTATTCTTT TTCTTACGTC TTCTGAGTAC TTCATACTTT CCAAATTTTT  
 430 440 450 460 470 480  
 CAACTGAAC TTTCTTCTTT TCTCATTGAA GTGGTTGGT TTTGGTCGCG TGATCAACGG  
 490 500 510 520 530 540  
 ATCCTACTTT TTTGAAACAA AATGTTTTG AAGTTTCACA GACTGATTTC GGGGTTTTTT  
 550 560 570 580 590 600  
 CAAAGAATAT ATTCCCTCTC GAGCAAGAGA AAATTCAGA AAATAGTAGT TTTTTCAAT  
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 670 680 690 700 710 720  
 GCTTCATACA AAAACATTAA ACTAGTGTGTT TTCCAGTTTT GTGTTCGTTT CATTTCCTCA  
 730 740 750 760 770 780  
 CCAAACGTGAC AATAATTACT TTCTGTGAAAC GTGTTTTGTA GGCAAGCTCC CGAATATTTT  
 790 800 810 820 830 840  
 TTTCTCTTCT CACGTCTTGT TATTTCTCG ATTTTATTTT CTGAATCTGT GCGGTTTCA  
 850 860 870 880 890 900  
 ATCAATTGAA TTGCGATAAT TATTCTATCA GAAATATATT TTCAGAAATC CAAATACCTCC  
 910 920 930 940 950 960  
 AGGTGCCAAT GCGGTGAAAG AAAATTTGA AGTTTATTCC TGAAATCACA CTACTCTTGC  
 970 980 990 1000 1010 1020  
 TTTTATTGTA ACACCTCTACA CAGGTTAGTT TCTCTTGAAT GTCCATTAA ATATTTATAG  
 1030 1040 1050 1060 1070 1080  
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 1090 1100 1110 1120 1130 1140  
 ATAACACTGCGT TTCTGGGTT TCTATCAGCA CTTACTAGCT GACAAAAACT TTTCCGTATT

1150 1160 1170 1180 1190 1200  
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1210 1220 1230 1240 1250 1260  
CTTTGATTA TATTGCTCGC ACCCTAAATG ACAGGTATTA GAAATTAACC GCTTTTCAGA  
1270 1280 1290 1300 1310 1320  
GTATTTTAA TCTTCTTAGT ACTAGTTAG TTCTTAAAT AAGAAACCAT CTAGTTTTC  
1330 1340 1350 1360 1370 1380  
ATTATCACTC AACTTCAGTC GGACAAATTT TAAATTTTT ACTCGATAAA AAAATTTAT  
1390 1400 1410  
AATTCAAGACA AATTATGTCT TCTCATTGTTT GATCGCT

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70	80	90	100	110	120
TCTCTTCCTG GATCCTTCAG AGCTCTTGTC AATTCTCAC GTTTTTTTT GTTTTTTCGT					
130	140	150	160	170	180
CGTTAATTG TGGAAACACA TATCCGTCT CTTGAAACA GCATCAGAAA ACTTTCTGCT					
190	200	210	220	230	240
CTCCGTGTCC TTCTACTTAC TCTGATTGCC TTAGTTAGTC ACATCGCAAG CAACAACATAA					
250	260	270	280	290	300
CTGCCAATGG GAGGAGCCAG TTGGAGCAGG GTGGGTGCTC GGTGCTCTTT TCAGAAGGTT					
310	320	330	340	350	360
TTCTCTTGTG CCAGCATGCT TTTTGAGGC TGTGTCATCA CAATGAACAT GTGTGAGTTC					
370	380	390	400	410	420
ATCCGTCTGG ATTATTCTTT TTCTTACGTC TTCTGAGTAC TTCATACTTT CCAAATTTTT					
430	440	450	460	470	480
CAAATGAACT TTTCTTCTTT TCTCATTGAA GTGGTTGGT TTTGGTCGCG TGATCAACGG					
490	500	510	520	530	540
ATCCTACTTT TTTGAAACAA AATGTTTTTG AAGTTTCACA GACTGATTTC GGGGTTTTTT					
550	560	570	580	590	600
CAAAGAATAT ATTCCCTCTC GAGCAAGAGA AAATTCCAGA AAATAGTAGT TTTTTCAAT					
610	620	630	640	650	660
TAGTCGTTTC ATTTGTACTA GCTAAAAAAC TTGCAACTTA TGGCTTTAAA ACATGTGTTG					
670	680	690	700	710	720
GCTTCATACA AAAACATTAA ACTAGTGTTC TTCCAGTTT GTGTTCGTTT CATTCTCA					
730	740	750	760	770	780
CCAAACTGAC AATAATTACT TTCTGTGAAC GTGTTTGTA GGCAAGCTCC CGAATATTTT					
790	800	810	820	830	840
TTTCTCTTCT CACGTCTTGT TATTTCTCG ATTTTATTTT CTGAATCTGT GCGGTTTCA					
850	860	870	880	890	900
ATCAATTGAA TTGCGATAAT TATTCTATCA GAAATATATT TTCAGAAATC CAAATACTCC					
910	920	930	940	950	960
AGGTGCCAAT GGGGTGAAAG AAAATTATGA AGTTTATTCC TGAAATCACA CTACTCTTGC					
970	980	990	1000	1010	1020
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1030	1040	1050	1060	1070	1080
TGCAAGGATA ATATAATTGA ATTGTTTTTG AGGAGTGCAA AGATTGAATA GTTTCTATA					
1090	1100	1110	1120	1130	1140
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1210	1220	1230	1240	1250	1260
TTAATATTGA GATTCATGTA CAGGTTGGTA GACTCTATAA TTGCACACCCA ATATGTGAAA					
1270	1280	1290	1300	1310	1320
GTTTCTTTA AAATTAAACT GCTGTAAATG ACTTTGAAT AAGTTTATCA GATAGAAATT					
1330	1340	1350	1360	1370	1380
GTCTGAACCTT TTTCGATTCAA ACTTTCCGAA CTTCAAAGCG GTTCCAAATT ACTCACTTCC					
1390	1400	1410	1420	1430	1440
ATTTATCTCT TTGCTACAAT TTCTCCCACA AAGCCTTTT CTTCATTAA CGTTCTTTT					
1450	1460	1470	1480	1490	1500
TATGTCGTTG TTCTTACAAA CAATTCGTC TCCTTGATGA ACTGCTTGAA CTGAGAATAG					
1510	1520	1530	1540	1550	1560
TCACATGAGG ATAAATTGTA TGGAATGACA AGTTTTGTGC CCAGAAGGCA GTTTGCAC					
1570	1580	1590	1600	1610	1620
GAACTTGTTG AGTTGCAGAC ACATCTCAAACACAGAAGA TGAGTGGAAA ACTAGTGAGA					
1630	1640	1650	1660	1670	1680
GACTGCCAAA AGTCGAAGGG ATAATGAAAA TTTGTTGCAA ATGAATTCTG CGAAGTTATG					
1690	1700	1710	1720	1730	1740
TGAAAAATTAA TTGGATTGGG ACTTGTGGGA GTGAAGAGAT GGGTCAAAAG CCATCAATCT					
1750	1760	1770	1780	1790	1800
TGAATGCTTC GGTCAAAGAT TTGTTTCTCA TATGTTTACA ACACTGAAAA CAATCTATCC					
1810	1820	1830	1840	1850	1860
TAGAAATGTT TGAACCACCC TCTAAAGTCC TTCCGTATAT TTTTCATCT TTATACCGAC					
1870	1880	1890	1900	1910	1920
CAGAATTCAA GAGTTGTTG AAATAACTTC CTCTTTTTG GAGAATATGT ACTCAGATTT					
1930	1940	1950	1960	1970	1980
TTACATTCAA AATTTATATA TTTTCAAATA GAAAAAGTGC CAAGTACCAAG AAACCTTTAT					
1990	2000	2010	2020	2030	2040
CAAGTTGGCG GCACTTTGGGA GAGTGAATTG GATGAAAAAG TGTTTGATAA GTTTGTCGGG					
2050	2060	2070	2080	2090	2100
CAAACCTGGTC CCCTGGGTGG GGAAATGGTG GCATTTTGG AAACATTTC ATAGTCGAAG					
2110	2120	2130	2140	2150	2160
AAGTGGAAACA AGAAATTGG AAAATAGAGA TACATATGTA TATGAAAATA GAATTGAACA					
2170	2180	2190	2200	2210	2220
CGAACTTATT TTTATTTCA GGATATGGGA AGCTTGAATG AAATATCATC CGACTTTGAA					
2230	2240	2250	2260	2270	2280
ATTGACATTT TATTCACCTCA ACTGTGGCAT GACTCGGCAC TTTCTTTGC TCATCTTCCG					

2290 2300 2310 2320 2330 2340  
 GCTTGTAAAGC GGTAAGAAAT CTTTGTATTA GAAGGGAAAAA ATATTTAAAT TAATGAATT  
 2350 2360 2370 2380 2390 2400  
 TCAGAAATAT CACAATGGAA ACACGACTTT TACCTAAGAT TTGGTCTCCA AACACGTGTA  
 2410 2420 2430 2440 2450 2460  
 TGATTAATTC AAAACGAACA ACCGTCGATG CATCACCATC GGAAAATGTG ATGGTTATTC  
 2470 2480 2490 2500 2510 2520  
 TGTACGAGGT ATGATTTTTG ATTTTGTGAC GTCACAAAACA GAGCATGTCT AAGGGCATGT  
 2530 2540 2550 2560 2570 2580  
 TGTAGCAAGA AAAAAACCGA TTCTTGTCTC TGTCGACGTT TCCTAAGTAT TGTGAATTAT  
 2590 2600 2610 2620 2630 2640  
 TTATAATACA TCACTCTAAT TACGTGAATA CTTACACCTT TAACTGGGTG AAGGATAAAA  
 2650 2660 2670 2680 2690 2700  
 TAGAGAAGGA GACGTTGAAA AAGCTCTTCG GTAGATTAAA GAGTCTAGAA TCGACATATG  
 2710 2720 2730 2740 2750 2760  
 TATTCAATGTT TCTCGGTTCA GGGAAATAAG TGATTTGGC GAAAAAGAGT TAGACGACAT  
 2770 2780 2790 2800 2810 2820  
 TTTTAGAAA ACTAAAACCA TATTCTCGAA CCCAAATCAG TCTAATGGTT TTCAGCAAA  
 2830 2840 2850 2860 2870 2880  
 AGTATGAAAT ATACAATGTT TGTTTCAGAA TACCCAGTAC AAAATTGAA GTTTTCAGA  
 2890 2900 2910 2920 2930 2940  
 ATGGAACAGT CTGGATTAAC CATCGTCTTA GTGTCAAATC ACCTTGCAAT TTGGATCTGC  
 2950 2960 2970 2980 2990 3000  
 GACAGTTCC TTTCGATACT CAAACTTGCA TATTAATCTT TGAATCCTAT AGTCATAACT  
 3010 3020 3030 3040 3050 3060  
 CAGAAGAAGT TGAACCTTCAT TGGATGGAAG AAGCTGTCAC ATTAATGAAG CCAATTCAAC  
 3070 3080 3090 3100 3110 3120  
 TTCCTGACTT TGATATGGTT CATTATTCAA CTAAAAAGGA AACTTTACTC TATCCAAACG  
 3130 3140 3150 3160 3170 3180  
 GCTACTGGGA TCAGCTCAA GTTACTTTCA CTTTCAAACG ACGATATGGA TTCTATATTA  
 3190 3200 3210 3220 3230 3240  
 TTCAAGCTA TGTTCCAACA TATCTTACAA TCATTGTATC TTGGGTTTCA TTCTGCATGG  
 3250 3260 3270 3280 3290 3300  
 AACCAAAAGC TCTGCCGGCA AGAACAACTG TCGGAATCTC ATCTCTTCTA GTTCTTACTT  
 3310 3320 3330 3340 3350 3360  
 TCCAGTTGG AAAATTTTG AAAATCTTC CAAGGGTTTC ATATGTGAAA GGTTTGTGTTT  
 3370 3380 3390 3400 3410 3420  
 TTTTCTTTTT CAACAAATA AAAAAAAGA TAAACAAATA TTTGTTTCAG CAATGGATGT

3410 3440 3450 3460 3470 3480  
 GTGGATGCTT GGATGCATAT CATTGTCCTT CGGAACCAGT GTAGAATTGG CATTGTTG  
 3490 3500 3510 3520 3530 3540  
 TTACATTTCC CGTTGTCAGA ACAGCGTAAG AAAGTGAGTT GGCATAAGAG TTTTCTCACG  
 3550 3560 3570 3580 3590 3600  
 TGGAGGGAAAG TAATTAAATT TTGGGTGTCA TATGAAAATA TCAAAAACAA TATCAGGAAA  
 3610 3620 3630 3640 3650 3660  
 TTGAATTCAGA CTATGATTC GTAGTAAACA AATTACAGCG CGGAACGACG ACGGAACGAA  
 3670 3680 3690 3700 3710 3720  
 ATGAGAAATT CTCAGGTGTG GGCAAACGGA TCGTGTAGAA CTAGAAGCAA CGGGTATGCA  
 3730 3740 3750 3760 3770 3780  
 AACGGGGGAT CTGTAATCTC ACATTATCAT CCAACAAGCA ATGGAAATGG GAATAATAAT  
 3790 3800 3810 3820 3830 3840  
 CGACATGATA CACCTCAAGT TACTGGAAGG TTAGCAATCT CTATGATAGC ATTTATCAAT  
 3850 3860 3870 3880 3890 3900  
 TATTAAGAA CTCTGGAATT AGTTTTAAA GTATAAATAA ATCTCTATT TTTGCGACCT  
 3910 3920 3930 3940 3950 3960  
 ACATTGAAC TAAATAGTTAT GTTTTACAGA GGATCACTTC ATCGAAACGG GCCACCATCT  
 3970 3980 3990 4000 4010 4020  
 CCATTAAACC TTCAAAATGAC TACATTTGAT TCGGAGATCC CTCTGACTTT TGATCAGGTG  
 4030 4040 4050 4060 4070 4080  
 AGTCTTACAT TGAGTTCAAA CTTTTGAAT TTAAGCGTTC TATCTGATAA AGTTCTTCGG  
 4090 4100 4110 4120 4130 4140  
 TGGTTTTATA ATTTTGATT CATAAACTTA CCCACTCCTT TCTCACTAAC ATTTTACCCCT  
 4150 4160 4170 4180 4190 4200  
 GTTCAGCTGC CAGTTCCAT GGAATCCGAT AGACCCCTGA TTGAAGAGGT AACTGTGAAA  
 4210 4220 4230 4240 4250 4260  
 GTAGTCAATT AATTCCCTGT GTTTCTACCC CACTCAATCC TTTTGTATT TTTGTTCACT  
 4270 4280 4290 4300 4310 4320  
 CTATCCACTA TCAATGTCTT ATCACCTCTA GATACTGTTT AGAAGAAAAT ATTGTTCAC  
 4330 4340 4350 4360 4370 4380  
 GTTATGGAAA TCACATATAC TTTGTTCTGG AATTGTATAT GTATGCTTTG AAAAAGCACA  
 4390 4400 4410 4420 4430 4440  
 TTAGAATACT ACAAACATTA GTTTCCATCA GATTTTGAT TTATCAAAC CGTTATATTA  
 4450 4460 4470 4480 4490 4500  
 GACACTCTTA AGTTATCATA TTCTAATTTC CAAGAATGTT ATATTTGAA GAAGCCGGTG  
 4510 4520 4530 4540 4550 4560  
 ATTGTCAAAA AGATTGAAAAA CTCCGAGTTT CTATATATGC GAAATTTCA CTTCAGCCCCA

45	4580	4590	4600	4610	4620
CACACACACA CACACATTCA CGAAACTTTG TGTGTTTAT GTTACTTATA TGTTATCTT					
4630	4640	4650	4660	4670	4680
TCTGTCGTAT CATGGTTTTC GGACTGAAAT TGTGTTAAC TGAAGTTATA TGTGAGCCAC					
4690	4700	4710	4720	4730	4740
ATTGATTAAA CCTCTGAGAG ATGCCCATTT GTACTCATTT TACGACTGTC TCATGTCCAA					
4750	4760	4770	4780	4790	4800
ACACCATGTT TATTGTAATT ACCAGGCTAC TATTTGCAGA TCGGATCAAC ATCACACCAC					
4810	4820	4830	4840	4850	4860
CCACCATCTG GATGTCCTGGC CAGATTCCAT CCCGAAGCAG TGGACAAATT CTCCATTGTA					
4870	4880	4890	4900	4910	4920
GCTTTCCAT TGGCATTAC AATGTTAAT GTTAGTTAAT CCACAGTTAA AAATTCCCAT					
4930	4940	4950	4960	4970	4980
AATCATAAAAT ATCTCGACTT TTCAGCTTGT CTACTGGTGG CACTATTTGT CTCAAACTTT					
4990	5000	5010	5020	5030	5040
CGATCAAAAC TATCAGTGAT TGAAGTTTAT CCCTTTAAT TCCAATAATT CACAGTTGCC					
5050	5060	5070	5080	5090	5100
GGTATCTACC TCCATTCTTT TCCGATGATT CGCAGTTTTT CACAGGGTTC AAATGTATCT					
5110	5120	5130	5140	5150	5160
CGTTCAATCT TTTTATGGTT ATTTCTCTTG AATGTCCATT TTAATATTTA TAGAACACTT					
5170	5180	5190	5200	5210	5220
TTATGTACAT TGTGTTGGTA TTCAATTGAA AAAACAAATGA AATTTATTC TAAATAACTG					
5230	5240	5250	5260	5270	5280
CGTTTCTGGG GTTTCTATCA GCACTTACTA GCTGACAAAA ACTTTCCGT ATTGGAATT					
5290	5300	5310	5320	5330	5340
AGATTTTAT GCAAGCAATG TTTCATTTC ACACAGTATA GTATTTATTC TTACTTTGA					
5350	5360	5370	5380	5390	5400
TTATATTGCT CGCACCCCTAA ATGACAGGTA TTAGAAATTA ACCGCTTTTC AGAGTATT					
5410	5420	5430	5440	5450	5460
TAATCTTCTT AGTACTAGTT TAGTTCTTTA AATAAGAAC CATCTAGTT TTCATTATCA					
5470	5480	5490	5500	5510	5520
CTCAACTTCA GTCGGACAAA TTTAAATT TTTACTCGAT AAAAAAATT TATAATTCA					
5530	5540	5550			
ACAAATTATG TCTTCTCATT TTTGATCGCT					

Fig. 7

1	20	30	40	50	60
ATGAAGTTA <u>TCCTGAAAT</u> CACACTACTC TTGCTTTAT <u>TATACACTC</u> TACACAGGCT					
70	80	90	100	110	120
AAAGGAAAAC GACGGAAATG TCCGGAGGGT GCGTGGTCGG AAGGAAAGAT TATGAACACG					
130	140	150	160	170	180
ATCATGAGCA ACTACACGAA <u>AATGTTGCC</u> GACCGGGAGG ACAGCGTACA AGTTAATATT					
190	200	210	220	230	240
GAGATTCATG TACAGGATAT GGGAAAGCTTG AATGAAATAT CATCCGACTT TGAAATTGAC					
250	260	270	280	290	300
ATTTTATTCA CTCAACTGTG GCATGACTCG GCACCTTCTT TTGCTCATCT TCCGGCTTGT					
310	320	330	340	350	360
AAGCGAAATA TCACAATGGA AACACGACTT TTACCTAAGA TTTGGTCTCC AAACACGTGT					
370	380	390	400	410	420
ATGATTAATT CAAAACGAAC AACCGTCCAT GCATCACCAT CGGAAAATGT GATGGTTATT					
430	440	450	460	470	480
CTGTACGAGA ATGGAACAGT CTGGATTAAC CATCGTCTTA GTGTCAAATC ACCTTGCAAT					
490	500	510	520	530	540
TTGGATCTGC GACAGTTCC TTTCGATACT <u>CAAACTTGCA</u> TATTAATCTT TGAATCCTAT					
550	560	570	580	590	600
AGTCATAACT CAGAAGAAGT TGAACATTCA TGGATGGAAG AAGCTGTCAC ATTAATGAAAG					
610	620	630	640	650	660
CCAATTCAAC TTCCTGACTT TGATATGGTT CATTATTCAA CTAAAAAGGA AACTTTACTC					
670	680	690	700	710	720
TATCCAAACG GGTACTGGGA TCAGCTTCAA GTTACTTTCA CTTTCAAACG ACGATATGGA					
730	740	750	760	770	780
TTCTATATTA TTCAAGCCTA TGTTCCAACA TATCTTACAA TCATTGTATC TTGGGTTTC					
790	800	810	820	830	840
TTCTGCATGG AACCAAAGC TCTGCCGGCA AGAACAACTG TCGGAATCTC ATCTCTTCTA					
850	860	870	880	890	900
<u>GT</u> TCTTACTT TCCAGTTGG AAATATTTG <u>AAAAATCTTC</u> CAAGGGTTTC ATATGTGAA					
910	920	930	940	950	960
GCAATGGATG TGTGGATGCT TGGATGCATA TCATTTGTCT TCGGAACCAT GGTAGAATTG					
970	980	990	1000	1010	1020
GCATTTGTTT GTTACATTTC CCGTTGTCAG AACAGCGTAA GAAACGCGGA ACGACGACGG					
1030	1040	1050	1060	1070	1080
GAACGAATGA GAAATTCTCA GGTGTGGGCA AACGGATCGT GTAGAACTAG AAGCAACGGG					
1090	1100	1110	1120	1130	1140
TATGCAAACG GGGGATCTGT AATCTCACAT TATCATCCAA CAAGCAATGG AAATGGGAAT					

Fig. 8  
Page 1

1150 1160 1170 1180 1190 1200  
AATAATCGAC ATGATACACC TCAAGTTACT GGAAGAGGAT CACTTCATCG AAACGGGCCA

1210 1220 1230 1240 1250 1260  
CCATCTCCAT TAAACCTTCA AATGACTACA TTTGATTCTGG AGATCCCTCT GACTTTGAT

1270 1280 1290 1300 1310 1320  
CAGCTGCCAG TTTCCATGGG ATCCGATAGA CCCCTGATTG AAGAGATGCG ATCAACATCA

1330 1340 1350 1360 1370 1380  
CCACCTCCAC CATCTGGATG TCTGGCCAGA TTCCATCCGG AAGCAGTGGA CAAATTCTCC

1390 1400 1410 1420 1430 1440  
ATTGTAGCTT TTCCATTGGC ATTTACAATG TTTAATCTTG TCTACTGGTG GCACTATTTG

1450 1460 1470  
TCTCAAACCTT TCGATCAAAA CTATCAGTGA

The MOD-1 Channel is Activated by Serotonin

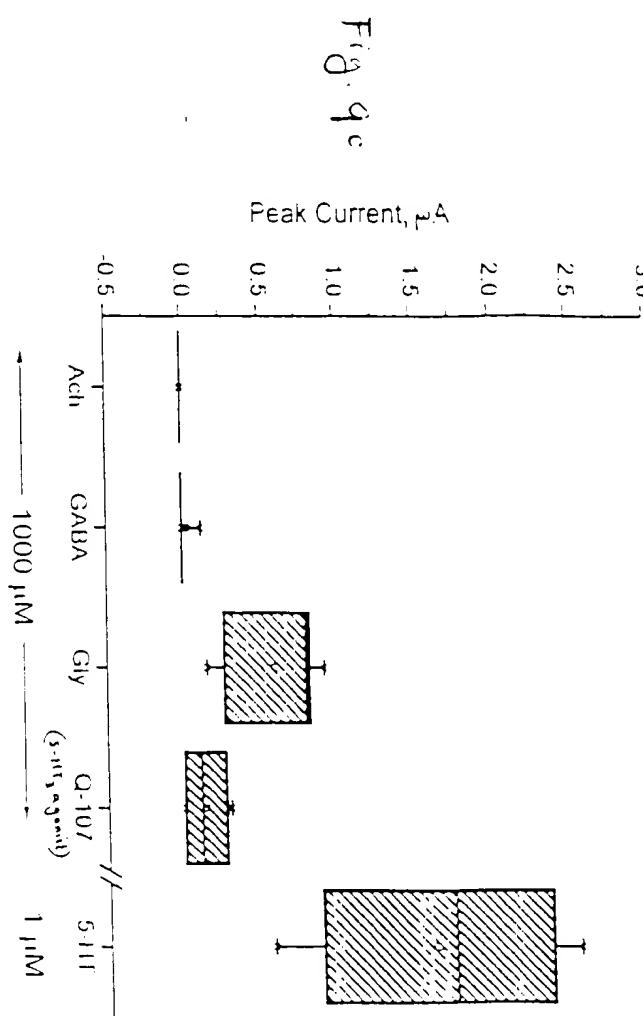
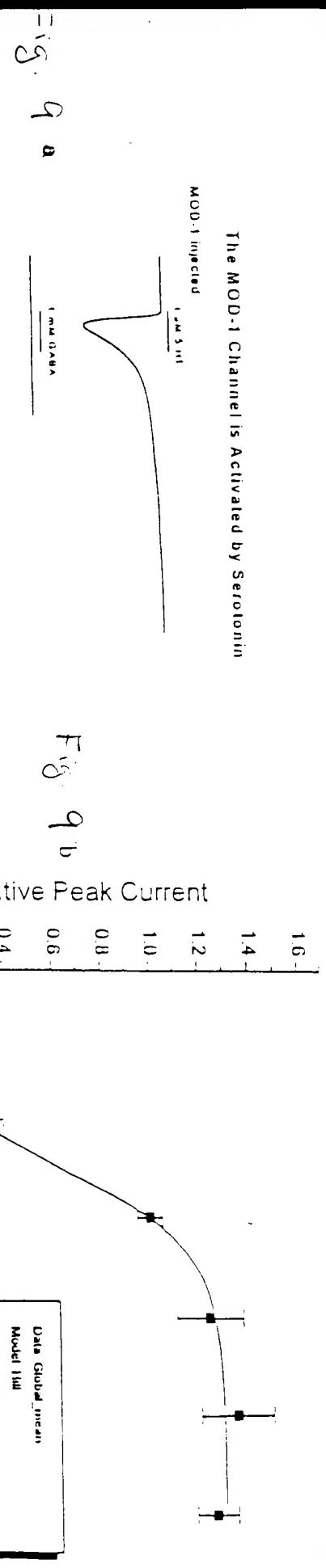


Fig. 9

# MOD-1 Selectivity

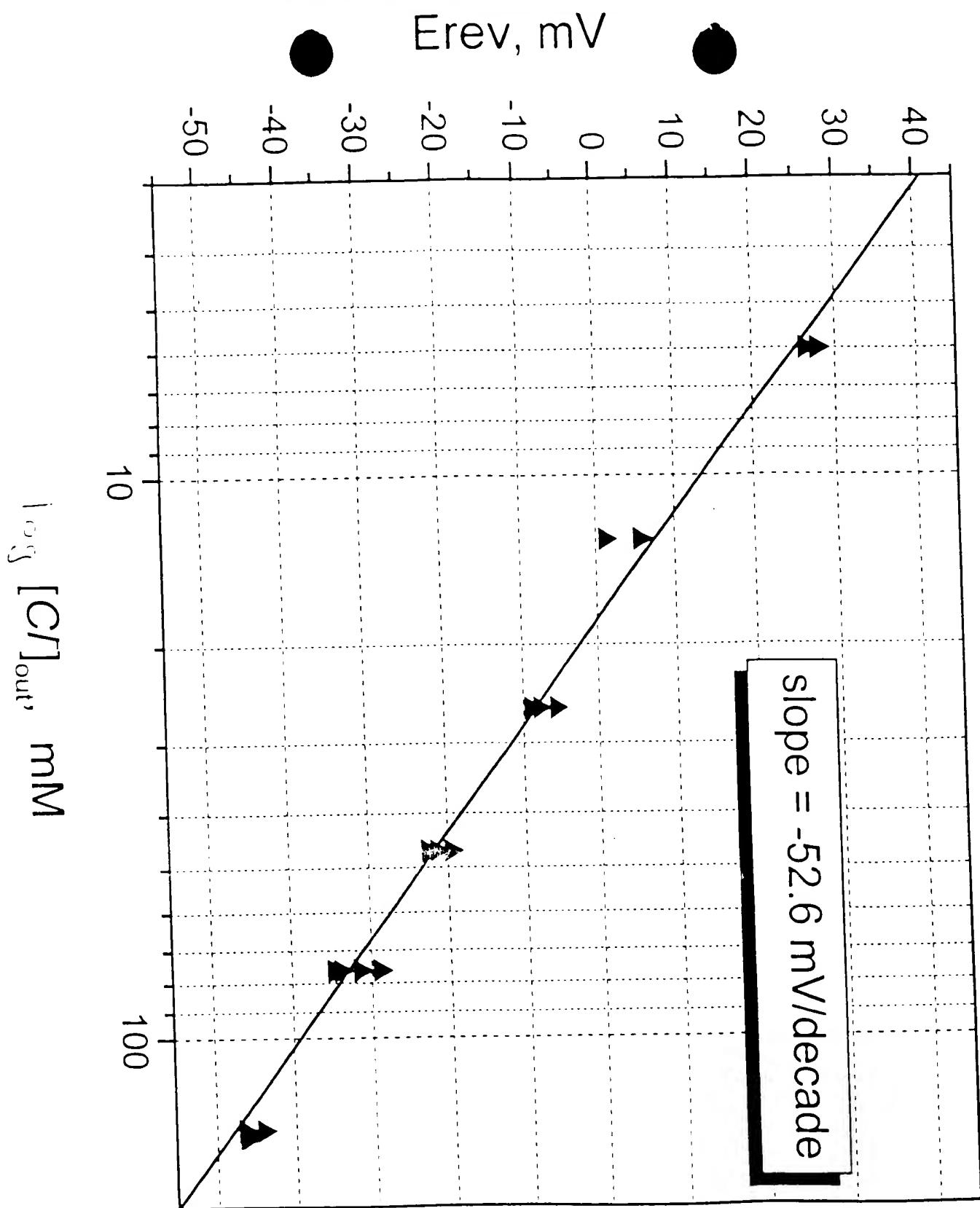


Fig. 10

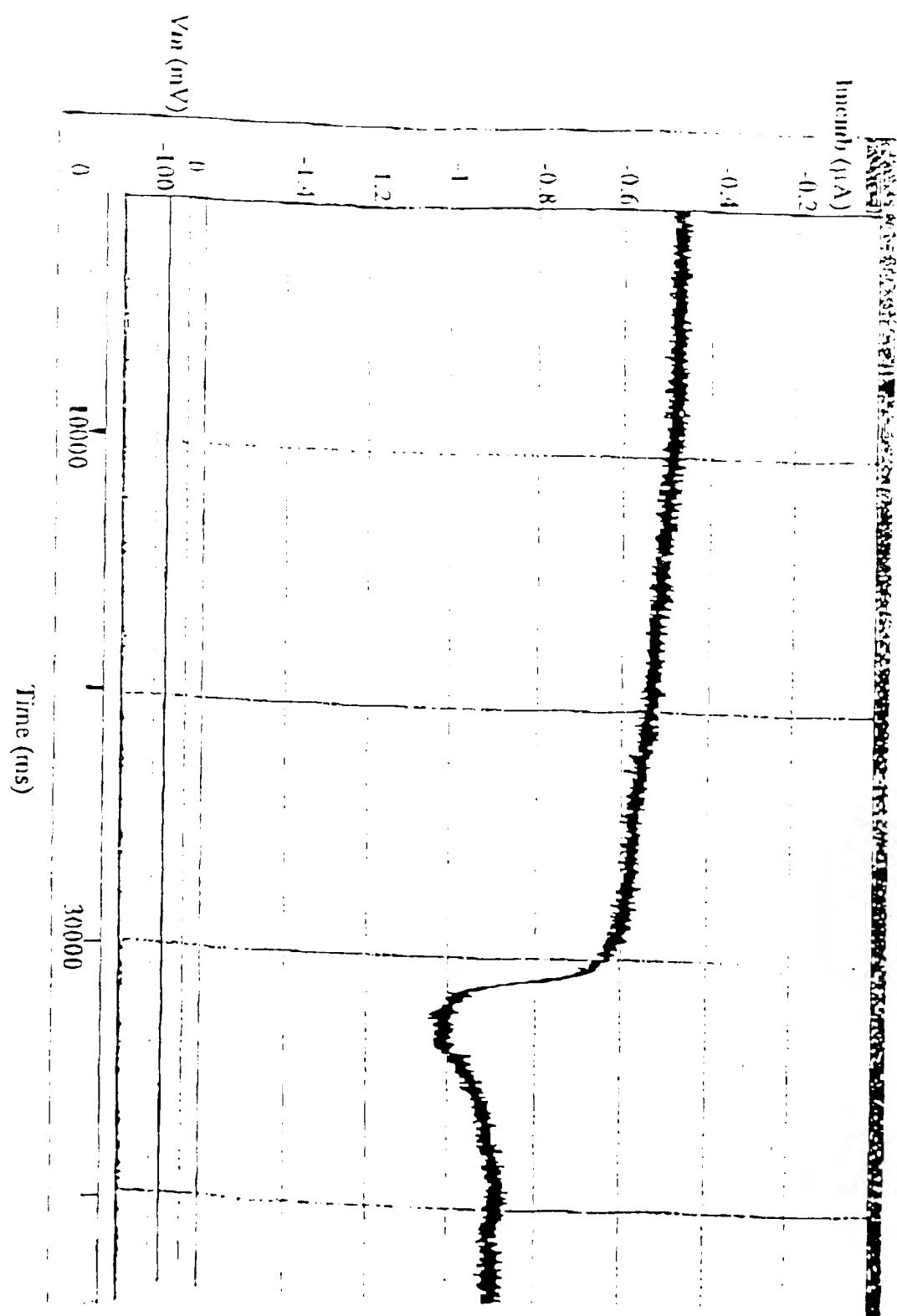


Figure 11

04/01/2010